AMENDMENTS TO THE CLAIMS

1-72. **(Canceled)**

- 73. **(Currently Amended)** A method of determining a definite quantity of target mRNA in a blood sample comprising:
 - (a) collecting whole blood;
 - (b) administering an anticoagulant to the whole blood;
 - (c) removing erythrocytes and blood components other than leukocytes from the whole blood to yield leukocytes;
 - (d) lysing the leukocytes with a lysis buffer containing spiked control RNA to produce a lysate comprising <u>target</u> mRNA and spiked control RNA, thereby obtaining amounts of <u>target</u> mRNA and spiked control RNA respectively, wherein said spiked control RNA is non-homologous to RNA from the whole blood sample;
 - (e) transferring the lysate to an oligo(dT)-immobilized plate to capture the mRNA;
 - (f) quantifying the <u>target</u> sample mRNA and <u>the</u> spiked control RNA, thereby obtaining values of <u>the target</u> sample mRNA and <u>the</u> spiked control RNA respectively;
 - (g) determining the percent recovery of <u>said</u> spiked control RNA by dividing the value of <u>said</u> spiked control RNA determined in step (f) by the amount of <u>said</u> spiked control RNA obtained in step (d); and
 - (h) determining the definite quantity of <u>target</u> mRNA by dividing the value of <u>said target</u> sample mRNA determined in step (f) by the percent recovery of <u>said</u> spiked control RNA determined in step (g).
 - 74. (Canceled)
- 75. **(Currently Amended)** The method of Claim 73, wherein step (b) (c) comprises filtration to yield leukocytes on a filter membrane.
 - 76. (Canceled)
- 77. **(Currently Amended)** The method of Claim 73, wherein <u>the anticoagulant</u> is heparin is administered to the whole blood prior to collection of leukocytes.
- 78. **(Currently Amended)** The method of Claim 73, wherein the whole blood is frozen and subsequently thawed prior to step (c) filtration.

- 79. **(Original)** The method of Claim 75, wherein the filter membrane is attached to a multi-well filter plate.
- 80. (Currently Amended) The method of Claim 79, wherein 10 to $1x10^{10}$ copies of spiked control RNA are applied to each <u>well of the multi-well</u> filter plate.
- 81. (Currently Amended) The method of Claim 79, wherein $1x10^5$ to $1x10^{10}$ copies of spiked control RNA are applied to each <u>well of the multi-well</u> filter plate.
- 82. **(Previously Presented)** The method of Claim 75, wherein the filter membrane is a polybutylene terephthalate (PBT) fibrous membrane.
- 83. (Currently Amended) The method of Claim 73 75, wherein the leukocytes are captured step (c) comprises filtration to yield leukocytes on a plurality of filter membranes layered together.
- 84. **(Original)** The method of Claim 75, additionally comprising washing the leukocytes on the filter membrane with hypotonic buffer to further remove erythrocytes and other blood components.
- 85. **(Original)** The method of Claim 84, additionally comprising drying the filter membrane.
- 86. **(Original)** The method of Claim 85, wherein the filter membrane is washed with ethanol and subjected to vacuum aspiration until the filter membrane is dry.
- 87. **(Original)** The method of Claim 73, wherein the immobilized plate comprises a multi-well oligo(dT)-immobilized plate.
- 88. (**Original**) The method of Claim 73, wherein the transfer of lysate to the oligo(dT)-immobilized plate comprises centrifugation.
- 89. **(Withdrawn)** The method of Claim 73, wherein the transfer of lysate to the oligo(dT)-immobilized plate comprises vacuum aspiration.
- 90. **(Withdrawn)** The method of Claim 73, wherein the transfer of lysate to the oligo(dT)-immobilized plate comprises applying positive pressure.
- 91. **(Currently Amended)** The method of Claim 73, wherein the quantification of <u>target</u> mRNA comprises cDNA synthesis of the <u>target</u> specific mRNA and amplification of <u>the</u> resulting cDNA.

92. **(Currently Amended)** The method of Claim 79 73, additionally comprising application of specific antisense primers to each well of the multi-well filter plate during said lysate transferring step.

93-214. (Canceled)

- 215. **(Currently Amended)** A method of high throughput quantification of a specific target mRNA, comprising the steps of:
 - (a) collecting whole blood;
 - (b) administering an anticoagulant to the whole blood;
 - (c) removing erythrocytes and blood components other than leukocytes from the whole blood by filtration to yield leukocytes on a filter membrane;
 - (d) lysing the leukocytes on said filter membrane with a lysis buffer comprising antisense primers specific to said specific target mRNA to produce a lysate comprising mRNA comprising said specific target mRNA with said antisense primers hybridized thereto;
 - (e) transferring the lysate to an oligo(dT)-immobilized plate to capture the specific target mRNA;
 - (f) removing non-hybridized materials from said oligo(dT)-immobilized plate;
 - (g) adding reverse transcriptase to said oligo(dT)-immobilized plate without addition of <u>further antisense</u> primers, thereby synthesizing cDNA formed by extension of both the immobilized oligo(dT) and the antisense primers,

wherein the cDNA formed by extension of <u>said</u> oligo(dT) remains immobilized to said plate, and the cDNA formed by extension of the antisense primers goes into solution as a result of displacement by the cDNA formed by extension of <u>said</u> oligo(dT) without heat denaturation <u>of said target mRNA and said cDNA formed by extension of the antisense primers;</u> and

(h) quantifying the specific target mRNA from by quantifying the cDNA in said cDNA solution.

216. (Canceled)

- 217. **(Currently Amended)** The method of Claim 215, wherein a plurality of different antisense primers for different specific target mRNAs are present in the lysis buffer.
- 218. (Currently Amended) The method of Claim 217, wherein each of said different mRNAs is amplified from the cDNA formed by extension of the antisense primers in step (g).
- 219. **(Previously Presented)** The method of Claim 217, wherein the cDNA solution is removed from the plate and the plate with the immobilized cDNA is stored for future use.
- 220. (Withdrawn) The method of Claim 73, wherein the mRNA quantified is β -actin mRNA.
- 221. (Withdrawn) The method of Claim 73, wherein the mRNA quantified is CD4 mRNA.
- 222. **(Withdrawn)** The method of Claim 73, wherein the mRNA of a translocation gene involved in leukemia is quantified.
- 223. **(Withdrawn)** The method of Claim 73, wherein the mRNA of cancer-specific genes from micrometastatic cancer is quantified.
- 224. **(Withdrawn)** The method of Claim 73, wherein virus-derived mRNA from infected white blood cells is quantified.
- 225. **(Withdrawn)** The method of Claim 224, wherein the quantified virus-derived mRNA is HIV.
- 226. **(Withdrawn)** The method of Claim 225, wherein the quantification of HIV mRNA is used to diagnose HIV.
- 227. **(Withdrawn)** The method of Claim 224, wherein the quantified virus-derived mRNA is CMV.
- 228. **(Withdrawn)** The method of Claim 227, wherein the quantification of virus-derived mRNA is used to diagnose CMV.
- 229. **(Withdrawn)** The method of Claim 224, wherein the quantification of virus-derived mRNA is used to monitor blood banks for the presence of viral diseases.
- 230. **(Withdrawn)** The method of Claim 224, wherein the quantification of virus-derived mRNA is used to study anti-viral drug sensitivity.

- 231. **(Currently Amended)** The method of Claim 73, wherein the <u>target mRNA</u> is mRNA of apoptosis genes involved in leukemia is quantified.
- 232. **(Currently Amended)** The method of Claim 73, wherein the <u>target mRNA</u> is mRNA of cytokines is quantified.
- 233. (Currently Amended) The method of Claim 73, wherein the <u>target mRNA</u> is <u>mRNA</u> responsible for apoptosis development, and wherein the quantification of mRNA is used to test the side effects of <u>anti-leukemia</u> <u>anti-cancer</u> drugs that induce mRNA responsible for apoptosis development in <u>leukocytes</u>.
- 234. **(Withdrawn)** The method of Claim 73, wherein the mRNA of DNA-repair genes is quantified.
- 235. **(Withdrawn)** The method of Claim 234, wherein the quantification of mRNA of DNA-repair genes is used to test the sensitivity of DNA-repair genes to radiation.
- 236. (Withdrawn) The method of Claim 73, wherein the mRNA of allergen response genes is quantified.
- 237. **(Withdrawn)** The method of Claim 236, wherein the quantification of mRNA of allergen response genes is used to test allergen stimulation.
 - 238. (Canceled)
 - 239. (Canceled)
- 240. **(New)** The method of Claim 217, wherein each of said different mRNAs is amplified from the cDNA formed by extension of the immobilized oligo(dT) in step (g).